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TCACCAGGTGCTCACCTTCCTCCTGCTCTTCGTGATCACCTCGGTGGCCTCTGAAAACGCCAGCACATCC
CGAGGCTGTGGGCTGGACCTCCTCCCTCAGTACGTGTCCCTGTGCGACCTGGACGCCATCTGGGGCATTTG
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GCCCTTCATCAAGGAGAAGGAGAAGAAGAGCCCTGTGGGCCTCCACTTCTGTTCCTCCTGGGGACCCCTG
GGCCTCTTTGGGCTGACGTTTGCCTTCATCATCCAGGAGGACGAGACCATCTGCTCTGTCCGCCGCTTCC
TCTGGGGCGTCTCTTTGCGCTCTGCTTCTCCTGCCTGCTGAGCCAGGCATGGCGCGTGGCGGAGGCTGGT
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TCTGTGCGGCAAGTTCAAGAGGTGGAAGCTGAACGGGGCCTTCTCCTCATCACAGCCTTCTCTCTGTG
CTCATCTGGGTGGCCTGGATGACCATGTACCTCTTCGGCAATGTCAAGCTGCAGCAGGGGGATGCCTGGA
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GATCCACTGCACCCTTCTGCCAGCCCTGCAGGAGAACACGCCCAACTACTTCGACACGTGCGAGCCCAGG
ATGCGGGAGACGGCCTTCGAGGAGGACGTGCAGCTGCCGCGGGCCTATATGGAGAACAAGGCCTTCTCCA
TGGATGAACACAATGCAGCTCTCCGAACAGCAGGATTTCCCAACGGCAGCTTGGGAAAAAGACCCAGTGG
CAGCTTGGGAAAAAGACCCAGCGCTCCGTTTAGAAGCAACGTGTATCAGCCAACTGAGATGGCCGTCGTG
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GTTCCAGAGAATCAGAATTTCTCTTACCGATTTGCCTCCCTGGCTGTGTCTTTCTTGAGGGAGAAATCGG
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GAAATGTGGCTGGGAAGACTGTTTCATCCTCTGGGGGTAGAACAGAACCAATTCACAGCTGGTGGGCC
AGACTGGTGTGTTGGTGGAGGTGGGGGGCTCCCACTCTTATCACCTCTCCCCAGCAAGTGCTGGACCCAG
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TTAACCTTTAAAGAAAAATGAAAAGGTTAGTGTTTGGGGGCCGGGGGAGGACTGACCGCTTCATAAGCC
AGTACGTCTGAGCTGAGTATGTTTCAATAAACCTTTTGATATTTCTCAAAAAAAAAAAAAAAAAAAAAA
(SEQ ID NO:1)

FIGURE 1A

MFVASERKMRAHQVLTFLLLFVITSVASENASTSRGCGLDLLPQYVSLCDLDAIWGIVVEAVAG
AGALITLLMLILLVRLPFIKEKEKKSPVGLHFLFLLGTLGLFGLTFAFIIQEDETICSVRRFL
WGVLFALCFSCLLSQAWRVRRLVRHGTGPAGWQLVGLALCLMLVQVIIAVEWLVLTVLRDTRPA
CAYEPMDFVMALIYDMVLLVVTGLGLALFTLCGKFKRWKLNGAFLITAFLSVLIWVAWMTMYLF
GNVKLQQGDAWNDPTLAITLAASGWVFVIFHAIPEIHCTLLPALQENTPNYFDTSQPRMRETA
EEDVQLPRAYMENKAFSMDEHNAALRTAGFPNGSLGKRPSGSLGKRPSAPFRSNVYQPTEMAVV
LNGGTIPTAPPSHTGRHLW (SEQ ID NO:2)

FIGURE 1B

underlined = deleted in targeting construct

bold = sequence flanking Neo insert in targeting construct

AGGTCGCAGGCGGGCGTGCGTGGAGCGGGGGCGGGCGCGCCGAGAGATGTGACTCG
 GGCCGAAGGCCAGCTGGAGCGTCGGCGCTGCGGGGCGGGGGT**CGAATGTTCTGGCA**
TCAGAGAGAAAGATGAGAGCTCACCAGGTGCTCACCTTCTCTGCTCTTCGTGATCACC
TCGGTGGCCTCTGAAAACGCCAGCACATCCCAGGCTGTGGGCTGGACCTCCTCCCTCAG
TACGTGTCCCTGTGCGACCTGGACGCCATCTGGGGCATTGTGGTGGAGGCGGTGGCCGGG
GCGGGCGCCCTGATCACACTGCTCCTGATGCTCATCCTCCTGGTGGGCTGCCCTTCATC
AAGGAGAAGGAGAAGAAGAGCCCTGTGGGCTCCACTTCTGTTCTCCTCCTGGGGACCCTG
GGCCTCTTTGGGCTGACGTTTGCCTTCATCATCCAGGAGGACGAGACCATCTGCTCTGTC
CGCCGCTTCTCTGGGGCGTCTCTTTGCGCTCTGCTTCTCCTGCCTGCTGAGCCAGGCA
TGGCGCGTGCGGAGGCTGGTGGCGCATGGCACGGGCCCCGCGGGCTGGCAGCTGGTGGGC
CTGGCGCTGTGCCTGATGCTGGTGCAAGTCATCATCGCTGTGGAGTGGCTGGTGTCTACC
GTGCTGCGTGACACAAGGCCAGCCTGCGCCTACGAGCCCATGGACTTTGTGATGGCCCTC
ATCTACGACATGGTACTGCTTGTGGTCACCCTGGGGCTGGCCCTCTTCACTCTGTGGCGC
AAGTTCAGAGGTGGAAGCTGAACGGGGCCTTCTCCTCATCACAGCCTTCTCTCTGTG
CTCATCTGGGTGGCCTGGATGACCATGTACCTCTTCGGCAATGTCAAGCTGCAGCAGGGG
GATGCCTGGAACGACCCACCTTGGCCATCACGCTGGCGGCCAGCGGCTGGGTCTTCGTC
ATCTTCCACGCCATCCCTGAGATCCACTGCACCCTTCTGCCAGCCCTGCAGGAGAACACG
CCCACTACTTCGACACGTCGACGCCAGGATGCGGGAGACGGCCTTCGAGGAGGACGTG
CAGCTGCCGCGGGCCTATATGGAGAACAAGGCCTTCTCCATGGATGAACACAATGCAGCT
CTCCGAACAGCAGGATTTCCCAACGGCAGCTTGGGAAAAAGACCCAGTGGCAGCTTGGGG
AAAAGACCCAGCGCTCCGTTTAGAAGCAACGTGTATCAGCCAAGTGAATGGCCGTCGTG
CTCAACGGTGGGACCATCCCAACTGCTCCGCCAAGTCACACAGGAAGACACCTTTGGTGA
AAGACTTTAAGTTCAGAGAATCAGAATTTCTCTTACCGATTTGCCTCCCTGGCTGTGTC
TTTCTTGAGGGAGAAATCGGTAACAGTTGCCGAACAGGCCGCTCACAGCCAGGAAATT
TGGAAATCCTAGCCAAGGGGATTTTCGTGTAAATGTGAACACTGACGAAGTAAAAGCTAA
CACCGACTGCCCCCCCCCTCCCTGCCACACACAGACACGTAATACCAGACCAACCTCA
ATCCCCGCAAACTAAAGCAAAAGCTAATTGCAAAATAGTATTAGGCTCACTGGAATGTGG
CTGGGAAGACTGTTTCATCCTCTGGGGGTAGAACAGAACCAATTCACAGCTGGTGGGCC
AGACTGGTGTGGTGGAGGTGGGGGGCTCCCACTCTTATCACCTCTCCCCAGCAAGTGC
TGGACCCCAAGTAGCCTCTTGGAGATGACCGTTGCGTTGAGGACAAATGGGGACTTTGCC
ACCGGCTTGCCCTGGTGGTTTGCACATTTACAGGGGGTTCAGGAGAGTTAAGGAGGTGTGG
GTGGGATTTCAAGGTGAGGCCCAACTGAATCGTGGGGTGAGCTTTATAGCCAGTAGAGGT
GGAGGGACCTGGCATGTGCCAAAGAAGAGGCCCTCTGGGTGATGAAGTGACCATCACAT
TTGGAAGTGATCAACCACTGTTCCTTCTATGGGGCTCTTGCTCTAATGTCTATGGTGAG
AACACAGGCCCCGCCCTTCCCTTGTAGAGCCATAGAAATATTCTGGCTTGGGGCAGCAG
TCCCTTCTTCCCTTGATCATCTCGCCCTGTTCCCTACACTTACGGGTGTATCTCCAAATCC
TCTCCCAATTTTATTCCTTATTTCATTTCAAGAGCTCCAATGGGGTCTCCAGCTGAAAGC
CCCTCCGGGAGGCAGGTGGGAAGGCAGGCACCACGGCAGGTTTTCGCGGATGATGTCACC
TAGCAGGGCTTCAGGGGTTCCTACTAGGATGCAGAGATGACCTCTCGCTGCCTCACAAAGC
AGTGACACCTCGGGTCCCTTCCGTTGCTATGGTGAAAAATTCCTGGATGGAATGGATCACA
TGAGGGTTTCTTGTGCTTTTGGAGGGTGTGGGGGATATTTGTTTGGTTTCTGTCAG
GTTCCATGAAAACAGCCCTTTTCCAAGCCCATGTTCCTGTGTCATGGTTTCCATCTGTCCCT
GAGCAAGTCATTCCTTTGTTATTTAGCATTTTCAACATCTCGGCCATTCAAAGCCCCAT
GTTCTCTGCACGTGTTTGGCCAGCATAACCTCTAGCATCGATTCAAAGCAGAGTTTAAACC
TGACGGCATGGAATGTATAAATGAGGGTGGGTCTTCTGCAGATACTCTAATCACTACAT
TGCTTTTCTATAAACTACCCATAAGCCTTTAACCTTTAAAGAAAAATGAAAAGGTTA
GTGTTTGGGGGCGGGGGAGGACTGACCGCTTCATAAGCCAGTACGCTGAGCTGAGTAT
GTTTCAATAAACCTTTTGATATTTCTCAAAAAAAAAAAAAAAAAAAAAA

FIGURE 2A

Gene Sequence
Structure *

526 bp

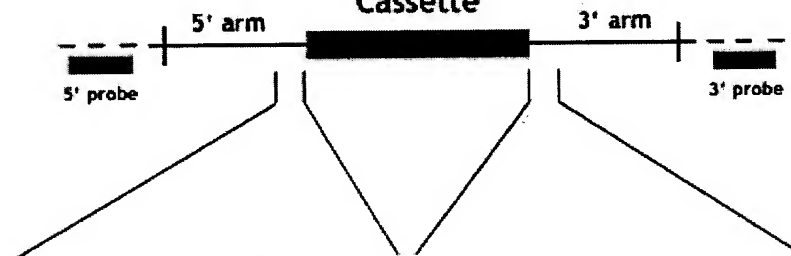
Sequence Deleted

594 bp

Size of full-length
cDNA: 2870 bp

Targeting Vector*
(genomic sequence)

LacZ-Neo
Cassette



Arm Length:
5': 4.2 kb
3': 0.7 kb

———— Targeting Vector
----- Endogenous Locus

* Not drawn to scale

5' >TGATGCTCATTCTCCTAGTGA
GACTACCC TTCATCAAGGACAAGG
AAAGGAAGCGGCTGTGTGCCTCC
ATTTCTCTTCTTGCTGGGGACCC
TGGGCCCTCTTTGGCCTGACGTTT
CCTTCATCATCCAGATGGACGAGA
CAATCTGCTCCATCCGACGCTTCC
TCTGGGGTGTCTCTTCGCGCTCT
GCTTTTCCGCT <3' (SEQ ID
NO: 3)

5' >GTGAGCCTGGCACTGTGCCTG
ATGCTGGTGCAGGTCATCATTGCC
ACTGAGTGGCTGGTGCTGACTGTG
CTGCGTGACACGAAGCCAGCCTGC
GCCTACGAGCCCATGGATTTTGTG
ATGGCGCTCATCTACGACATGGTG
CTGCTGGCCATCACCTGGCCAG
TCCCTCTTACGCTGTGTGGCAAG
TTCAAACGGTG <3' (SEQ ID
NO: 4)

FIGURE 2B

Hot Plate Test

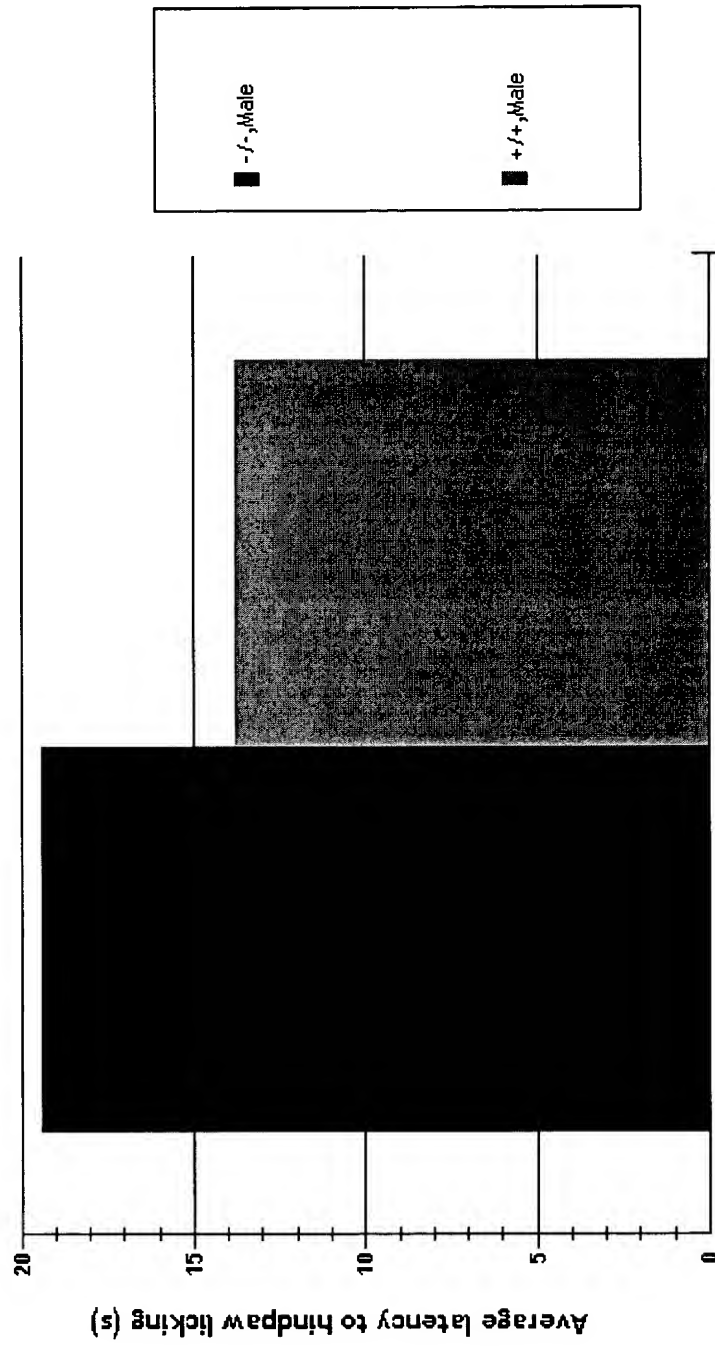


FIGURE 3